#2

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RAW SEQUENCE LISTING DATE: 05/23/2001
PATENT APPLICATION: US/09/851,614 TIME: 13:07:33

Input Set : A:\Seqlist.txt

Output Set: C:\CRF3\05232001\I851614.raw

```
3 <110> APPLICANT: Deo, Yashwant M.
              Keler, Tibor
       <120> TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO DENDRITIC
      9 <130> FILE REFERENCE: MXI-166
C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/851,614
                                                                  ENTERED
C--> 11 <141> CURRENT FILING DATE: 2001-05-08
     11 <150> PRIOR APPLICATION NUMBER: USSN 60/203,126
     12 <151> PRIOR FILING DATE: 2000-05-08
     14 <150> PRIOR APPLICATION NUMBER: USSN 60/230,739
     15 <151> PRIOR FILING DATE: 2000-09-07
     17 <160> NUMBER OF SEQ ID NOS: 7
     19 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     21 <210> SEQ ID NO: 1
     22 <211> LENGTH: 321
     23 <212> TYPE: DNA
     24 <213> ORGANISM: Homo sapiens
     26 <220> FEATURE:
     27 <221> NAME/KEY: CDS
     28 <222> LOCATION: (1)...(321)
     30 <400> SEQUENCE: 1
     31 gac atc cag atg acc cag tet eca tec tea etg tet gea tet gta gga
     32 Asp Ile Gln Met Thr Gln Ser Pro-Ser Ser Leu Ser Ala Ser Val Gly
     35 gac aga gtc acc atc act tgt cgg gcg agt cag ggt att agc agg tgg
     36 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Ser Arg Trp
                                       · · 25
                     20
     39 tta god tgg tat dag dag aaa dda gag aaa god oot aag too dtg atd
     40 Leu Ala Trp Tyr Gln Gln Lys Pro Glu Lys Ala Pro Lys Ser Leu Ile
     41
                 35
                                     40
                                                          45
     43 tat get gea tee agt ttg caa agt ggg gte eea tea agg tte age gge
                                                                           192
     44 Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly
                                 55
                                                                           240
     47 agt gga tet ggg aca gat tte act ete ace ate age gge etg cag eet
     48 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Gly Leu Gln Pro
                             70
     51 gaa gat ttt gca act tat tac tgc caa cag tat aat agt tac cct cgg
                                                                           288
     52 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Tyr Asn Ser Tyr Pro Arg
     53
                                                                           321
     55 acg ttc ggc caa ggg acc aag gtg gaa atc aaa 1
     56 Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys
                    100
                                        105
     60 <210> SEQ ID NO: 2
     61 <211> LENGTH: 107
     62 <212> TYPE: PRT
     63 <213> ORGANISM: Homo sapiens
     65 <400> SEQUENCE: 2
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## (

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Input Set : A:\Seqlist.txt

Output Set: C:\CRF3\05232001\I851614.raw

66 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Vai Gly 67 1 5 10 15	
68 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Ser Arg Trp 69 20 25 30	
70 Leu Ala Trp Tyr Gln Gln Lys Pro Glu Lys Ala Pro Lys Ser Leu Ile 71 35 40 45	
72 Tyr Ala Ala Ser Ser Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly 73 50 55 60	
74 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Gly Leu Gln Pro 75 65 70 75 80	
76 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Tyr Asn Ser Tyr Pro Arg 77 85 90 95	
78 Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys 79 100 105	
82 <210> SEQ ID NO: 3 83 <211> LENGTH: 348	
84 <212> TYPE: DNA	
85 <213> ORGANISM: Homo sapiens	
87 <220> FEATURE:	
88 <221> NAME/KEY: CDS	
89 <222> LOCATION: (1)(348) 91 <400> SEQUENCE: 3	
92 gag gtg cag ctg gtg cag tot gga gca gag gtg aaa aag coo ggg gag	48
93 Glu Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Glu	
94 1 5 10 15	
96 tot otg agg ato too tgt aag ggt tot gga gac agt ttt acc acc tac	96
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Tyr	96
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98 20 25 30	
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Tyr	96
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Tyr 98 20 25 30 100 tgg atc ggc tgg gtg cgc cag atg ccc ggg aaa ggc ctg gag tgg atg	
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98 20 25 30 100 tgg atc ggc tgg gtg cgc cag atg ccc ggg aaa ggc ctg gag tgg atg 101 Trp Ile Gly Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Trp Met 102 35 40 45 104 ggg atc atc tat cct ggt gac tct gat acc ata tac agc ccg tcc ttc	
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98 20 25 30 100 tgg atc ggc tgg gtg cgc cag atg ccc ggg aaa ggc ctg gag tgg atg 101 Trp Ile Gly Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Trp Met 102 35 40 45 104 ggg atc atc tat cct ggt gac tct gat acc ata tac agc ccg tcc ttc 105 Gly Ile Ile Tyr Pro Gly Asp Ser Asp Thr Ile Tyr Ser Pro Ser Phe	144
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98 20 25 30 100 tgg atc ggc tgg gtg cgc cag atg ccc ggg aaa ggc ctg gag tgg atg 101 Trp Ile Gly Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Trp Met 102 35 40 45 104 ggg atc atc tat cct ggt gac tct gat acc ata tac agc ccg tcc ttc 105 Gly Ile Ile Tyr Pro Gly Asp Ser Asp Thr Ile Tyr Ser Pro Ser Phe 106 50 55 60	144 192
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98 20 25 30  100 tgg atc ggc tgg gtg cgc cag atg ccc ggg aaa ggc ctg gag tgg atg 101 Trp Ile Gly Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Trp Met 102 35 40 45  104 ggg atc atc tat cct ggt gac tct gat acc ata tac agc ccg tcc ttc 105 Gly Ile Ile Tyr Pro Gly Asp Ser Asp Thr Ile Tyr Ser Pro Ser Phe 106 50 55 60  108 caa ggc cag gtc acc atc tca gcc gac aag tcc atc agc acc gcc tac	144
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98 20 25 30 100 tgg atc ggc tgg gtg cgc cag atg ccc ggg aaa ggc ctg gag tgg atg 101 Trp Ile Gly Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Trp Met 102 35 40 45 104 ggg atc atc tat cct ggt gac tct gat acc ata tac agc ccg tcc ttc 105 Gly Ile Ile Tyr Pro Gly Asp Ser Asp Thr Ile Tyr Ser Pro Ser Phe 106 50 55 60	144 192
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98 20 25 30  100 tgg atc ggc tgg gtg cgc cag atg ccc ggg aaa ggc ctg gag tgg atg 101 Trp Ile Gly Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Trp Met 102 35 40 45  104 ggg atc atc tat cct ggt gac tct gat acc ata tac agc ccg tcc ttc 105 Gly Ile Ile Tyr Pro Gly Asp Ser Asp Thr Ile Tyr Ser Pro Ser Phe 106 50 55 60  108 caa ggc cag gtc acc atc tca gcc gac aag tcc atc agc acc gcc tac 109 Gln Gly Gln Val Thr Ile Ser Ala Asp Lys Ser Ile Ser Thr Ala Tyr 110 65 70 75 80  112 ctg cag tgg agc agc ctg aag gcc tcg gac acc gcc atg tat tac tgt	144 192
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98 20 25 30  100 tgg atc ggc tgg gtg cgc cag atg ccc ggg aaa ggc ctg gag tgg atg 101 Trp Ile Gly Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Trp Met 102 35 40 45  104 ggg atc atc tat cct ggt gac tct gat acc ata tac agc ccg tcc ttc 105 Gly Ile Ile Tyr Pro Gly Asp Ser Asp Thr Ile Tyr Ser Pro Ser Phe 106 50 55 60  108 caa ggc cag gtc acc atc tca gcc gac aag tcc atc agc acc gcc tac 109 Gln Gly Gln Val Thr Ile Ser Ala Asp Lys Ser Ile Ser Thr Ala Tyr 110 65 70 75 80  112 ctg cag tgg agc agc ctg aag gcc tcg gac acc gcc atg tat tac tgt 113 Leu Gln Trp Ser Ser Leu Lys Ala Ser Asp Thr Ala Met Tyr Tyr Cys	144 192 240
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98 20 25 30  100 tgg atc ggc tgg gtg cgc cag atg ccc ggg aaa ggc ctg gag tgg atg 101 Trp Ile Gly Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Trp Met 102 35 40 45  104 ggg atc atc tat cct ggt gac tct gat acc ata tac agc ccg tcc ttc 105 Gly Ile Ile Tyr Pro Gly Asp Ser Asp Thr Ile Tyr Ser Pro Ser Phe 106 50 55 60  108 caa ggc cag gtc acc atc tca gcc gac aag tcc atc agc acc gcc tac 109 Gln Gly Gln Val Thr Ile Ser Ala Asp Lys Ser Ile Ser Thr Ala Tyr 110 65 70 75 80  112 ctg cag tgg agc agc ctg aag gcc tcg gac acc gcc atg tat tac tgt 113 Leu Gln Trp Ser Ser Leu Lys Ala Ser Asp Thr Ala Met Tyr Tyr Cys 114 85 90 95	144 192 240 288
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98	144 192 240
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98	144 192 240 288
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98	144 192 240 288
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98 20 25 30 100 tgg atc ggc tgg gtg cgc cag atg ccc ggg aaa ggc ctg gag tgg atg 101 Trp Ile Gly Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Trp Met 102 35 40 45 104 ggg atc atc tat cct ggt gac tct gat acc ata tac agc ccg tcc ttc 105 Gly Ile Ile Tyr Pro Gly Asp Ser Asp Thr Ile Tyr Ser Pro Ser Phe 106 50 55 60 108 caa ggc cag gtc acc atc tca gcc gac aag tcc atc agc acc gcc tac 109 Gln Gly Gln Val Thr Ile Ser Ala Asp Lys Ser Ile Ser Thr Ala Tyr 110 65 70 75 80 112 ctg cag tgg agc agc ctg aag gcc tcg gac acc gcc atg tat tac tgt 113 Leu Gln Trp Ser Ser Leu Lys Ala Ser Asp Thr Ala Met Tyr Tyr Cys 85 90 95 116 acg aga ggg gac cgg ggc gtt gac tac tgg ggc cag gga acc ctg gtc 117 Thr Arg Gly Asp Arg Gly Val Asp Tyr Trp Gly Gln Gly Thr Leu Val 118 100 105 110 120 acc gtc tcc tca 121 Thr Val Ser Ser	144 192 240 288 336
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98	144 192 240 288 336
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98	144 192 240 288 336
97 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr 98	144 192 240 288 336





## RAW SEQUENCE LISTING DATE: 05/23/2001 PATENT APPLICATION: US/09/851,614 TIME: 13:07:33

Input Set : A:\Seqlist.txt

Output Set: C:\CRF3\05232001\I851614.raw

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128 <213> ORGANISM: Homo sapiens
     130 <400> SEQUENCE: 4
     131 Glu Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Glu
     133 Ser Leu Arg Ile Ser Cys Lys Gly Ser Gly Asp Ser Phe Thr Thr Tyr
                                         25
     135 Trp Ile Gly Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Trp Met
     137 Gly Ile Ile Tyr Pro Gly Asp Ser Asp Thr Ile Tyr Ser Pro Ser Phe
                                 55
     139 Gln Gly Gln Val Thr Ile Ser Ala Asp Lys Ser Ile Ser Thr Ala Tyr
     141 Leu Gln Trp Ser Ser Leu Lys Ala Ser Asp Thr Ala Met Tyr Tyr Cys
     143 Thr Arg Gly Asp Arg Gly Val Asp Tyr Trp Gly Gln Gly Thr Leu Val
                     100
                                         105
     145 Thr Val Ser Ser
     146
                115
    149 <210> SEQ ID NO: 5
     150 <211> LENGTH: 15
     151 <212> TYPE: PRT
     152 <213> ORGANISM: Homo sapiens
     154 <220> FEATURE:
     155 <221> NAME/KEY: VARIANT
     156 <222> LOCATION: (1)...(15) /
     157 <223> OTHER INFORMATION: Xaa = Any Amino Acid
     159 <400> SEQUENCE: 5
W--> 160 Asp Asp Xaa Xaa Gln Phe Leu Ile Xaa Xaa Glu Asp Xaa Lys Arg
     161 1
                          5
     164 <210> SEQ ID NO: 6
     165 <211> LENGTH: 15
     166 <212> TYPE: PRT
     167 <213> ORGANISM: Homo sapiens
    169 <400> SEQUENCE: 6
     170 Leu Asp Thr Arg Gln Phe Leu Ile Tyr Asn Glu Asp His Lys Arg
    171 1
    174 <210> SEQ ID NO: 7
     175 <211> LENGTH: 20
     176 <212> TYPE: PRT
    177 <213> ORGANISM: Homo sapiens
     179 <400> SEQUENCE: 7
     180 Leu Leu Asp Thr Arg Gln Phe Leu Ile Tyr Leu Glu Asp Thr Lys Arg
     181 1
     182 Cys Val Asp Ala
     183
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## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/851,614

DATE: 05/23/2001 TIME: 13:07:34

Input Set : A:\Seqlist.txt

Output Set: C:\CRF3\05232001\I851614.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:160 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5